

## Sequence Listing

### SEQUENCE LISTING

<110> Steinkasserer, Alexander

<120> Use of Soluble Forms of CD83 and Nucleic Acids Encoding them for the Treatment or Prevention of Diseases

<130> 032723woJH

<140>

<141>

<160> 12

<170> PatentIn ver. 2.1

<210> 1

<211> 618

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(615)

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gct ccc gcg acg ccg gag gtg aag gtg gct tgc tcc gaa gat gtg gac	96
Ala Pro Ala Thr Pro Glu Val Lys Val Ala Cys Ser Glu Asp Val Asp	
20 25 30	
ttg ccc tgc acc gcc ccc tgg gat ccg cag gtt ccc tac acg gtc tcc	144
Leu Pro Cys Thr Ala Pro Trp Asp Pro Gln Val Pro Tyr Thr Val Ser	
35 40 45	
tgg gtc aag tta ttg gag ggt ggt gaa gag agg atg gag aca ccc cag	192
Trp Val Lys Leu Leu Glu Gly Glu Glu Arg Met Glu Thr Pro Gln	
50 55 60	
gaa gac cac ctc agg gga cag cac tat cat cag aag ggg caa aat ggt	240
Glu Asp His Leu Arg Gly Gln His Tyr His Gln Lys Gly Gln Asn Gly	
65 70 75 80	
tct ttc gac gcc ccc aat gaa agg ccc tat tcc ctg aag atc cga aac	288
Ser Phe Asp Ala Pro Asn Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn	
85 90 95	
act acc agc tgc aac tcg ggg aca tac agg tgc act ctg cag gac ccg	336
Thr Thr Ser Cys Asn Ser Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro	
100 105 110	
gat ggg cag aga aac cta agt ggc aag gtg atc ttg aga gtg aca gga	384
Asp Gly Gln Arg Asn Leu Ser Gly Lys Val Ile Leu Arg Val Thr Gly	
115 120 125	
tgc cct gca cag cgt aaa gaa gag act ttt aag aaa tac aga gcg gag	432
Cys Pro Ala Gln Arg Lys Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu	
130 135 140	
att gtc ctg ctg ctg gct ctg gtt att ttc tac tta aca ctc atc att	480
Ile Val Leu Leu Leu Ala Leu Val Ile Phe Tyr Leu Thr Leu Ile Ile	
145 150 155 160	
ttc act tgt aag ttt gca cgg cta cag agt atc ttc cca gat ttt tct	528

Sequence Listing

Phe	Thr	Cys	Lys	Phe	Ala	Arg	Leu	Gln	Ser	Ile	Phe	Pro	Asp	Phe	Ser	
165																175
aaa	gct	ggc	atg	gaa	cga	gct	ttt	ctc	cca	gtt	acc	tcc	cca	aat	aag	576
Lys	Ala	Gly	Met	Glu	Arg	Ala	Phe	Leu	Pro	Val	Thr	Ser	Pro	Asn	Lys	
180								185							190	
cat	tta	ggg	cta	gtg	act	cct	cac	aag	aca	gaa	ctg	gtt	tga			618
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195							200								205	

<210> 2  
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<212> PRT  
<213> Homo sapiens

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Ala	Pro	Ala	Thr	Pro	Glu	Val	Lys	Val	Ala	Cys	Ser	Glu	Asp	Val	Asp	
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Leu	Pro	Cys	Thr	Ala	Pro	Trp	Asp	Pro	Gln	Val	Pro	Tyr	Thr	Val	Ser	
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Trp	Val	Lys	Leu	Leu	Glu	Gly	Gly	Glu	Glu	Arg	Met	Glu	Thr	Pro	Gln	
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Glu	Asp	His	Leu	Arg	Gly	Gln	His	Tyr	His	Gln	Lys	Gly	Gln	Asn	Gly	
				65		70			75					80		
Ser	Phe	Asp	Ala	Pro	Asn	Glu	Arg	Pro	Tyr	Ser	Leu	Lys	Ile	Arg	Asn	
				85			90						95			
Thr	Thr	Ser	Cys	Asn	Ser	Gly	Thr	Tyr	Arg	Cys	Thr	Leu	Gln	Asp	Pro	
				100				105					110			
Asp	Gly	Gln	Arg	Asn	Leu	Ser	Gly	Lys	Val	Ile	Leu	Arg	Val	Thr	Gly	
				115				120				125				
Cys	Pro	Ala	Gln	Arg	Lys	Glu	Glu	Thr	Phe	Lys	Lys	Tyr	Arg	Ala	Glu	
				130			135					140				
Ile	Val	Leu	Leu	Leu	Ala	Leu	Val	Ile	Phe	Tyr	Leu	Thr	Leu	Ile	Ile	
					145		150			155				160		
Phe	Thr	Cys	Lys	Phe	Ala	Arg	Leu	Gln	Ser	Ile	Phe	Pro	Asp	Phe	Ser	
				165				170					175			
Lys	Ala	Gly	Met	Glu	Arg	Ala	Phe	Leu	Pro	Val	Thr	Ser	Pro	Asn	Lys	
				180				185				190				
His	Leu	Gly	Leu	Val	Thr	Pro	His	Lys	Thr	Glu	Leu	Val				
				195				200				205				

<210> 3  
<211> 2051  
<212> DNA  
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<220>

Sequence Listing

<221> CDS

<222> (14)..(601)

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Sequence Listing

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gtcatctaca agctatggtg agatgcaggt gaagcagggt catggaaat ttgaacactc 931  
tgagctggcc ctgtgacaga ctcctgagga cagctgtcct ctcctacatc tggatacat 991  
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catttatttt ttaatcttc atgtacttgt caaagaagaa ttttcatgt ttttcaaag 1291  
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taatgagctg ggcccttcc tcatttgctt cccaaagaga ttttgcctca ctaatggtgt 1951  
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<210> 4  
<211> 196  
<212> PRT  
<213> Mus musculus

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Ala Asp Leu Pro Cys Thr Ala Pro Trp Asp Pro Gln Leu Ser Tyr Ala  
35 40 45

Val Ser Trp Ala Lys Val Ser Glu Ser Gly Thr Glu Ser Val Glu Leu  
50 55 60

Pro Glu Ser Lys Gln Asn Ser Ser Phe Glu Ala Pro Arg Arg Arg Ala  
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65	70	75	80
Tyr Ser Leu Thr Ile Gln Asn Thr Thr Ile Cys Ser Ser Gly Thr Tyr			
85	90	95	
Arg Cys Ala Leu Gln Glu Leu Gly Gly Gln Arg Asn Leu Ser Gly Thr			
100	105	110	
Val Val Leu Lys Val Thr Gly Cys Pro Lys Glu Ala Thr Glu Ser Thr			
115	120	125	
Phe Arg Lys Tyr Arg Ala Glu Ala Val Leu Leu Phe Ser Leu Val Val			
130	135	140	
Phe Tyr Leu Thr Leu Ile Ile Phe Thr Cys Lys Phe Ala Arg Leu Gln			
145	150	155	160
Ser Ile Phe Pro Asp Ile Ser Lys Pro Gly Thr Glu Gln Ala Phe Leu			
165	170	175	
Pro Val Thr Ser Pro Ser Lys His Leu Gly Pro Val Thr Leu Pro Lys			
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Thr Glu Thr val			
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<210> 5  
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<212> DNA  
<213> Artificial Sequence  
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<223> Description of Artificial Sequence: primer for CD83ext

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<212> DNA  
<213> Artificial Sequence  
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<223> Description of Artificial Sequence: primer for CD83ext

<400> 6  
aattagaatt ctcaaatctc cgctctgtat t 31

<210> 7  
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<212> DNA  
<213> Artificial Sequence  
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<223> Description of Artificial Sequence: partial sequence of pGEX2ThCD83ext  
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<222> (28)..(417)

## Sequence Listing

<210> 8

<211> 139

<212> PRT

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<223> Description of Artificial Sequence: partial sequence of pGEX2ThCD83ext

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Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro  
10 15 20

Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu  
25 30 35

Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly  
40 45 50 55

Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn  
60 65 70

Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser  
75 80 85

Sequence Listing

Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu  
 90 95 100

Ser Gly Lys Val Ile Leu Arg Val Thr Gly Cys Pro Ala Gln Arg Lys  
 105 110 115

Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile  
 120 125 130

<210> 9

<211> 435

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: partial sequence of pGEX2ThCD83ext\_mut129\_CtoS

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<221> CDS

<222> (1)..(417)

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<221> mat\_peptide

<222> (28)..(417)

<400> 9

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gtg aag gtg gct tgc tcc gaa gat gtg gac ttg ccc tgc acc acc gcc ccc	96
Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro	
10                    15            20	

tgg gat ccg cag gtt ccc tac acg gtc tcc tgg gtc aag tta ttg gag	144
Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu	
25                    30            35	

ggt ggt gaa gag agg atg gag aca ccc cag gaa gac cac ctc agg gga	192
Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly	
40                    45            50                55	

cag cac tat cat cag aag ggg caa aat ggt tct ttc gac gcc ccc aat	240
Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn	
60                    65            70	

gaa agg ccc tat tcc ctg aag atc cga aac act acc agc tgc aac tcg	288
Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser	
75                    80            85	

ggg aca tac agg tgc act ctg cag gac ccg gat ggg cag aga aac cta	336
Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu	
90                    95            100	

agt ggc aag gtg atc ttg aga gtg aca gga tcc cct gca cag cgt aaa	384
Ser Gly Lys Val Ile Leu Arg Val Thr Gly Ser Pro Ala Gln Arg Lys	
105                    110            115	

gaa gag act ttt aag aaa tac aga gcg gag att tgagaattca tcgtgact	435
Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile	
120                    125            130	

Sequence Listing

<210> 10  
<211> 139  
<212> PRT  
<213> Artificial Sequence  
<223> Description of Artificial Sequence: partial sequence of pGEX2ThCD83ext\_mut129\_Ctos

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Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu  
-5 -1 1 5

Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro  
10 15 20

Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu  
25 30 35

Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly  
40 45 50 55

Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn  
60 65 70

Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser  
75 80 85

Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu  
90 95 100

Ser Gly Lys Val Ile Leu Arg Val Thr Gly Ser Pro Ala Gln Arg Lys  
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Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile  
120 125 130

<210> 11  
<211> 32  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Description of Artificial Sequence: primer sense-pGEX2ThCD83

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<210> 12  
<211> 66  
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<223> Description of Artificial Sequence: primer antisense-CD83extra\_mutantCtos

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ggggat 66

<210> 13

Sequence Listing

<211> 209

<212> PRT

<213> Homo sapiens

<400> 13

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1 5 10 15

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20 25 30

Glu Asp Val Asp Leu Pro Cys Thr Ala Pro Trp Asp Pro Gln Val Pro  
35 40 45

Tyr Thr Val Ser Trp Val Lys Leu Leu Glu Gly Gly Glu Glu Arg Met  
50 55 60

Glu Thr Pro Gln Glu Asp His Leu Arg Gly Gln His Tyr His Gln Lys  
65 70 75 80

Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn Glu Arg Pro Tyr Ser Leu  
85 90 95

Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser Gly Thr Tyr Arg Cys Thr  
100 105 110

Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu Ser Gly Lys Val Ile Leu  
115 120 125

Arg Val Thr Gly Cys Pro Ala Gln Arg Lys Glu Glu Thr Phe Lys Lys  
130 135 140

Arg Arg Ala Glu Ile Val Leu Leu Leu Ala Leu Val Ile Phe Tyr Leu  
145 150 155 160

Thr Leu Ile Ile Phe Thr Cys Lys Phe Ala Arg Leu Gln Ser Ile Phe  
165 170 175

Pro Asp Phe Ser Lys Ala Gly Met Glu Arg Ala Phe Leu Pro Val Thr  
180 185 190

Ser Pro Asn Lys His Leu Gly Leu Val Thr Pro His Lys Thr Glu Leu  
195 200 205

Val  
209